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TEM & TRUE QUENCE LISTING
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QUENCE LISTING <110> Ryazanov, Alexey <120> MAMMALIAN ALPHA-KINASE PROTEINS, NUCLEIC ACIDS AND DIAGNOSTIC AND THERAPEUTIC USES THEREOF <130> 601-1-098CIP 09/832,292 <140> <141> 2001-04-10 <150> 09/632,131 2001-08-03 <151> <160> 45 <170> PatentIn version 3.1 <210> 1 <211> 238 <212> PRT <213> Homo sapiens <400> Gly Glu Trp Leu Asp Asp Glu Val Leu Ile Lys Met Ala Ser Gln Pro 10 Phe Gly Arg Gly Ala Met Arg Glu Cys Phe Arg Thr Lys Lys Leu Ser 20 25 Asn Phe Leu His Ala Gln Gln Trp Lys Gly Ala Ser Asn Tyr Val Ala Lys Arg Tyr Ile Glu Pro Val Asn Arg Asp Val Tyr Phe Glu Asp Val Arg Leu Gln Met Glu Ala Lys Leu Trp Gly Asp Asp Tyr Asn Arg His Lys Pro Pro Lys Gln Val Asp Ile Met Gln Met Cys Ile Ile Glu Leu Lys Asp Arg Pro Gly Lys Pro Leu Phe His Leu Asp His Tyr Ile Asp Gly Lys Tyr Ile Lys Tyr Asn Ser Asn Ser Gly Phe Val Arg Asp Asp Asn Ile Arg Leu Thr Pro Gln Ala Phe Ser His Phe Thr Phe Glu Arg Ser Gly His Gln Leu Ile Val Val Asp Ile Gln Gly Val Gly Asp Leu 150

Tyr Thr Asp Pro Gln Ile His Thr Glu Thr Gly Thr Asp Phe Gly Asn 165 170 175

Gly Asn Leu Gly Val Arg Gly Met Ala Leu Phe Phe Tyr Ser His Ala 180 185 190

Cys Asn Arg Ile Cys Glu Ser Met Gly Leu Ala Pro Phe Asp Leu Ser 195 200 205

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Leu Thr Pro Gln Ala Phe Ser His Phe Thr Phe Glu Arg Ser Gly His 130 135 140

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Met Thr Gln Leu Lys Asn Gly Thr Lys Phe Val Leu Lys Leu Tyr Lys 65 70 75 80

Lys Glu Ala Glu Gln Gln Ala Ser Arg Glu Leu Tyr Phe Glu Asp Val $85 \hspace{1cm} 90 \hspace{1cm} 95$

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- Lys Ser Glu Arg Tyr Ser Ser Ser Gly Ser Pro Ala Asn Ser Phe His 65 70 75 80
- Phe Lys Glu Ala Asn Lys His Ala Ile Gln Lys Ala Lys His Met Pro 85 90 95
- Asp Pro Trp Ala Glu Phe His Leu Glu Asp Ile Ala Thr Glu Arg Ala 100 105 110
- Thr Arg His Arg Tyr Asn Ala Val Thr Gly Glu Trp Leu Asp Asp Glu 115 120 125
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Met Leu Phe Thr Gly Gly Tyr Gly Leu Glu Lys Asp Pro Gln Arg Ser 675 680 685

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Lys Thr Glu Leu Cys Gly Ser Thr Gly Ser Pro Ala Ser Ser Phe His

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Thr Arg His Arg Tyr Asn Ala Val Thr Gly Glu Trp Leu Lys Asp Glu 115 120 125

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Ile Met Gln Met Cys Ile Ile Glu Leu Lys Asp Arg Pro Gly Gln Pro 210 215 220

Leu Phe His Leu Glu His Tyr Ile Glu Gly Lys Tyr Ile Lys Tyr Asn 225 230 235 240

Ser Asn Ser Gly Phe Val Arg Asp Asp Asn Ile Arg Leu Thr Pro Gln 245 250 255

Ala Phe Ser His Phe Thr Phe Glu Arg Ser Gly His Gln Leu Ile Val 260 265 270

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Thr Glu Lys Gly Thr Asp Phe Gly Asp Gly Asn Leu Gly Val Arg Gly 290 295 300

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Met Gly Leu Thr Pro Phe Asp Leu Ser Pro Arg Glu Gln Asp Ala Val 325 330 335

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His Phe Thr Glu Gln Met Glu Asp Gln Leu Glu Lys Thr Met Lys Val 100 105 110

Val Arg Asn His Thr Asp Ser Leu Gly Gly Asn Val Gln Thr Lys Leu 115 120 125

Asp Glu Gly Ile Glu Lys Cys Met Ala Phe Ala Lys Lys Val Glu Gln 130 135 140

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Lys Asp Asn Arg Ser Glu Leu Glu Gly Leu Glu Lys Asp Cys Lys Asn 405 410 415

385

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Lys Lys Ser Asp Ser Leu Leu Leu Met Gln Asn Asn Leu Lys Lys 435 440 445

Tyr Asn Glu Phe Val Asp Arg Glu Arg Asp Arg Glu Ser Glu Arg Leu 450 460

Lys Leu Gln Asp Ser Ile Lys Arg Leu Glu Gln Asn Gln Lys Lys Ile 465 470 475 480

Glu Ala Glu Ile Gln Glu Gly Asn Glu Gln Val Glu Arg Val Leu Arg 485 490 495

Glu Glu Ala Ser Ile Ser Pro Ile Ser Ser Val Pro Lys Ser Pro Ile 500 505 510

Thr Thr Lys Arg Ser Ser Ile Ile Leu Asn Ser Pro Pro Met Thr Ser 515 520 525

Gln Gln Ser Ser Pro Lys Ile Gln Asp Leu Leu Ser Ser Gly Ser 530 535 540

Ser Ser Val Ser Gly Ile Asn Ile Ser Ser Glu Thr Gly Glu Met Gly 545 550 560

Ile Leu Trp Glu Phe Asp Pro Ile Ile Asn Lys Trp Ile Arg Leu Ser 565 570 575

Met Lys Leu Lys Val Glu Arg Lys Pro Phe Ala Glu Gly Ala Leu Arg 580 585 590

Glu Ala Tyr His Thr Val Ser Leu Gly Val Gly Thr Asp Glu Asn Tyr 595 600 605

Pro Leu Gly Thr Thr Lys Leu Phe Pro Pro Ile Glu Met Ile Ser 610 615 620

Pro Ile Ser Lys Asn Asn Glu Ala Met Thr Gln Leu Lys Asn Gly Thr 625 630 635 640

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- Leu Met Ser Trp Val Val Glu Leu Ile Asp Arg Ser Pro Ser Ser Asn 690 695 700
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- Pro Gln Ala Phe Ser His Phe Thr Tyr Glu Leu Ser Asn Lys Gln Met 740 745 750
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- Pro Glu Leu Met Pro Ser Asp Asn Thr Ile Lys Val Gly Ala Lys Gln 835 840 845
- Leu Pro Lys Ala Glu Phe Ser Lys Lys Asp Leu Lys Cys Val Ser Thr 850 860
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- Lys Ser Ile Glu Ser Ile Ala Cys Asn Ser Asn Tyr Ile Phe Thr Ser 915 920 925
- Ser Pro Asp Asn Thr Ile Lys Val His Ile Ile Arg Ser Gly Asn Thr 930 935 940
- Lys Cys Ile Glu Thr Leu Val Gly His Thr Gly Glu Val Asn Cys Val 945 950 955 960
- Val Ala Asn Glu Lys Tyr Leu Phe Ser Cys Ser Tyr Asp Lys Thr Ile 965 970 975
- Lys Val Trp Asp Leu Ser Thr Phe Lys Glu Ile Lys Ser Phe Glu Gly 980 985 990
- Val His Thr Lys Tyr Ile Lys Thr Leu Ala Leu Ser Gly Arg Tyr Leu 995 1000 1005
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- Thr Leu Ser Met Leu Phe Asn Met Gln Gly His Glu Asp Trp Val 1025 1030 1035
- Leu Ser Leu His Cys Thr Ala Ser Tyr Leu Phe Ser Thr Ser Lys 1040 1045
- Asp Asn Val Ile Lys Ile Trp Asp Leu Ser Asn Phe Ser Cys Ile 1055 1060 1065
- Asp Thr Leu Lys Gly His Trp Asn Ser Val Ser Ser Cys Val Val 1070 1080
- Lys Asp Arg Tyr Leu Tyr Ser Gly Ser Glu Asp Asn Ser Ile Lys 1085 1090 1095
- Ser His Ser Leu Gly Val Lys Cys Leu Met Val Phe Asn Asn Gln

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<213> Dictyostelium discoideum

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<211> 732

<212> PRT

<213> Dictyostelium discoideum

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Pro Phe Ala Glu Gln Ala Phe Arg Lys Ala Tyr His Thr Leu Asp Leu 145 150 155 160

Ser Lys Ser Gly Ala Ser Gly Arg Tyr Val Ser Lys Ile Gly Lys Lys 165 170 175

Pro Thr Pro Arg Pro Ser Tyr Phe Glu Asp Val Lys Met Gln Met Ile 180 185 190

Ala Lys Lys Trp Ala Asp Lys Tyr Asn Ser Phe Lys Pro Pro Lys Lys 195 200 205

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Ser Asp Leu Ile Cys Gly Ala Glu Pro Tyr Val Glu Gly Gln Tyr Arg 225 230 235 240

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Pro Gln Ser Phe Ser His Phe Thr Tyr Glu His Ser Asn His Gln Leu 260 265 270

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Lys Gly Phe Glu Lys Phe Leu Asp Thr His Lys Cys Asn Ala Ile Cys 305 310 315 320

Gln Tyr Leu Asn Leu Gln Ser Ile Asn Pro Lys Ser Glu Lys Ser Asp 325 330 335

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Gln Tyr Leu Phe Ser Gly Ser Asn Asp Lys Ile Thr Lys Val Trp Asp 595 600 605

Leu Lys Thr Phe Arg Cys Asn Tyr Thr Leu Lys Gly His Thr Lys Trp 610 615 620

Val Thr Thr Ile Cys Ile Leu Gly Thr Asn Leu Tyr Ser Gly Ser Tyr 625 630 635 640

Asp Lys Thr Ile Arg Val Trp Asn Leu Lys Ser Leu Glu Cys Ser Ala 645 650 655

Thr Leu Arg Gly His Asp Arg Trp Val Glu His Met Val Ile Cys Asp 660 665 670

Lys Leu Leu Phe Thr Ala Ser Asp Asp Asn Thr Ile Lys Ile Trp Asp 675 680 685

Leu Glu Thr Leu Arg Cys Asn Thr Thr Leu Glu Gly His Asn Ala Thr 690 695 700

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<212> DNA

<213> C. elegans

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<213> C. elegans

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Asn Asp Val Val Ile Glu Lys Pro Arg Met Asp Pro Leu His Val Arg 50 60

Lys Leu Met Glu Thr Trp Arg Lys Ala Ala Arg Arg Ala Arg Thr Asn 70 75 80

Tyr Ile Asp Pro Trp Lys Glu Phe Asn Ile His Glu Tyr Pro Val Gln 85 90 95

Arg Ala Lys Arg Tyr Arg Tyr Ser Ala Ile Arg Lys Gln Trp Thr Glu 100 105 110

Asp Ile Val Asp Val Arg Leu His Pro Asp Ser Phe Ala Arg Gly Ala 115 120 125

Met Arg Glu Cys Tyr Arg Leu Lys Lys Cys Ser Lys His Gly Thr Ser 130 135 140

Gln Asp Trp Ser Ser Asn Tyr Val Ala Lys Arg Tyr Ile Cys Gln Val 145 150 155 160

Asp Arg Val Leu Phe Asp Asp Val Arg Leu Gln Met Asp Ala Lys 165 170 175

Leu Trp Ala Glu Glu Tyr Asn Arg Tyr Asn Pro Pro Lys Lys Ile Asp 180 185 190

The Val Gln Met Cys Val Ile Glu Met Ile Asp Val Lys Gly Ser Pro 195 200 205

Leu Tyr His Leu Glu His Phe Ile Glu Gly Lys Tyr Ile Lys Tyr Asn 210 215 220

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Phe Ser His Phe Thr Phe Glu Arg Ser Gly His Gln Met Met Val Val 245 250 255

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Val Ala Met Glu Val Ala Ala Lys Gln Lys Lys Ser Cys Ile Val Pro 325 330 335

Pro Thr Val Phe Glu Ala Arg Arg Asn Arg Ile Ser Ser Glu Cys Val 340 345 350

His Val Glu His Gly Ile Ser Met Asp Gln Leu Arg Lys Arg Lys Thr 355 360 365

Leu Asn Gln Ser Ser Thr Asp Leu Ser Ala Lys Ser His Asn Glu Asp 370 375 380

Cys Val Cys Pro Glu Cys Ile Pro Val Val Glu Gln Leu Cys Glu Pro 385 390 395 400

Cys Ser Glu Asp Glu Glu Asp Glu Glu Asp Tyr Pro Arg Ser Glu 405 410 415

Lys Ser Gly Asn Ser Gln Lys Ser Arg Arg Ser Arg Met Ser Ile Ser 420 425 430

Thr Arg Ser Ser Gly Asp Glu Ser Ala Ser Arg Pro Arg Lys Cys Gly 435 440 445

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n Gl
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Asp Glu Asp Val Pro Gln Val Thr Gly His Gln Phe Ser Val Leu Gly 515 520 525

Gln Ile His Ile Asp Leu Ser Arg Tyr His Glu Leu Gly Arg Phe Val530 . 535 540

Glu Val Asp Ser Glu His Lys Glu Met Leu Glu Gly Ser Glu Asn Asp 545 550 555 560

Ala Arg Val Pro Ile Lys Tyr Asp Lys Gln Ser Ala Ile Phe His Leu 565 570 575

Asp Ile Ala Arg Lys Cys Gly Ile Leu Glu Ala Val Leu Thr Ser Ala 580 585 590

His Ile Val Leu Gly Leu Pro His Glu Leu Leu Lys Glu Val Thr Val 595 600 605

Asp Asp Leu Phe Pro Asn Gly Phe Gly Glu Glu Asn Gly Ile Arg 610 615 620

Ala Asp Lys Gly Gln Lys Pro Cys Asp Leu Glu Glu Phe Gly Ser Asp 625 630 635 640

Leu Met Glu Ile Ala Ala Glu Met Gly Asp Lys Gly Ala Met Leu Tyr $645 \hspace{1.5cm} 650 \hspace{1.5cm} 655$

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Thr Asp Tyr Lys Lys Ser Ile Asp Trp Tyr Gln Arg Val Val Gly Phe 675 680 685

Gln Glu Glu Leu Asp Ser Asp Cys Gly Lys Thr Thr Phe Ser Ser 690 700

Phe Ala Pro Leu Thr Arg His Glu Ile Leu Ala Lys Met Ala Glu Met 705 710 715 720

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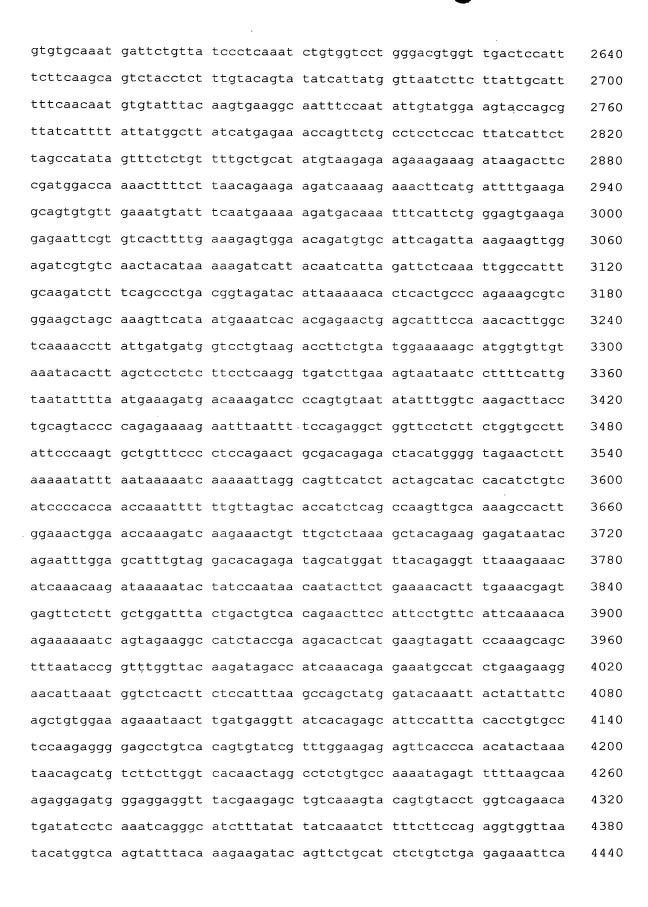
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Phe Gln Gly Ser His Ser Tyr Arg Ala Lys Tyr Val Arg Leu Ser 100 105 110

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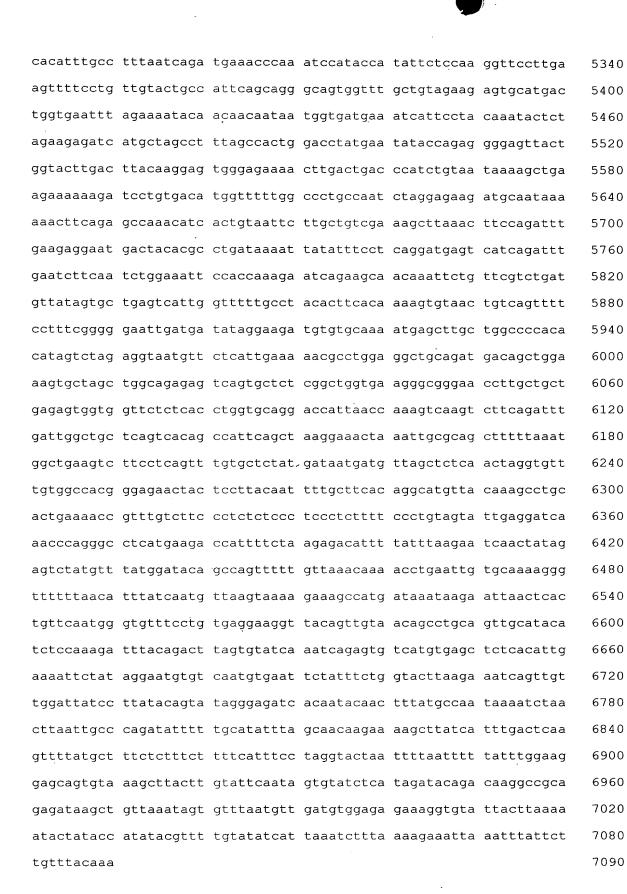
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Lys Gln His Ala Cys Phe Thr Ala Ser Leu Ala Met Lys Tyr Ser Asp 50 60

Val Arg Leu Gly Glu His Phe Asn Gln Ala Ile Glu Glu Trp Ser Val 65 70 75 80

Glu Lys His Thr Glu Gln Ser Pro Thr Asp Ala Tyr Gly Val IIe Asn 85 90 95

Phe Gln Gly Ser His Ser Tyr Arg Ala Lys Tyr Val Arg Leu Ser 100 105 110

Tyr Asp Thr Lys Pro Glu Ile Ile Leu Gln Leu Leu Leu Lys Glu Trp 115 120 125

Gln Met Glu Leu Pro Lys Leu Val Ile Ser Val His Gly Gly Met Gln 130 135 140

Lys Phe Glu Leu His Pro Arg Ile Lys Gln Leu Leu Gly Lys Gly Leu 145 150 155 160

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Asn Thr Gly Val Ala Lys His Val Gly Asp Ala Leu Lys Glu His Ala 180 185 190

Ser Arg Ser Ser Arg Lys Ile Cys Thr Ile Gly Ile Ala Pro Trp Gly 195 200 205

Val Ile Glu Asn Arg Asn Asp Leu Val Gly Arg Asp Val Val Ala Pro 210 215 220

Cys Leu Lys Leu Ala Val Ser Ser Arg Leu Arg Pro Phe Val Ala His 725 730 735

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Met Trp Tyr Tyr Ser Asp Gln Asn Ala Ser Ser Ser Lys Glu Ser Ala 770

Ser Val Lys Glu Tyr Asp Leu Glu Arg Gly His Asp Glu Lys Leu Asp 795

Glu Asn Gln His Phe Gly Leu Glu Ser Gly His Gln His Leu Pro Trp 805

Thr Arg Lys Val Tyr Glu Phe Tyr Ser Ala Pro Ile Val Lys Phe Trp 820 825 830

Phe Tyr Thr Met Ala Tyr Leu Ala Phe Leu Met Leu Phe Thr Tyr Thr 835 840 845

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- Ser Ile Tyr Ile Phe Thr Asn Ala Ile Glu Val Val Arg Glu Val Ser 865 870 875 880
- Ile Ser Glu Pro Gly Lys Phe Thr Gln Lys Val Lys Val Trp Ile Ser 885 890 895
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- Leu Ile Tyr Cys Ile Asp Ile Ile Phe Trp Phe Ser Arg Leu Leu Asp 930 935 940
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- Lys Met Thr Ala Asn Met Phe Tyr Ile Val Ile Ile Met Ala Ile Val 965 970 975
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- Tyr His Glu Lys Pro Trp Leu Pro Pro Pro Leu Ile Leu Leu Ser

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- Tyr Ser Ala Ile Glu Arg Asn Asn Leu Met Arg Leu Ser Gln Thr 1700 1710
- Ile Pro $\,$ Phe Thr Pro Val Gln Leu Phe Ala Gly Glu Glu Ile Thr $\,$ 1715 $\,$ $\,$ 1720 $\,$ 1725
- Val Tyr Arg Leu Glu Glu Ser Ser Pro Leu Asn Leu Asp Lys Ser 1730 1740
- Met Ser Ser Trp Ser Gln Arg Gly Arg Ala Ala Met Ile Gln Val 1745 1750
- Leu Ser Arg Glu Glu Met Asp Gly Gly Leu Arg Lys Ala Met Arg 1760 1765 1770
- Val Val Ser Thr Trp Ser Glu Asp Asp Ile Leu Lys Pro Gly Gln

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Thr Thr Gln Ala Ser Asp His Leu Arg Gln Pro Gln Glu Asn Arg Asp

Lys Thr Pro Ile Trp Asn Ser Gly Ser Thr Ser Leu Ser Arg Ser Phe 70 75

Leu Thr Arg Ser Pro Asn Glu Val His Lys Ile Ser Thr Ser Leu Lys 90

Ser Pro Gln Glu Pro His His Tyr Ser Ala Ile Glu Arg Asn Asn 100 105

Leu Met Arg Leu Ser Gln Thr Ile Pro Phe Thr Pro Ile Gln Leu Phe 115 120 125

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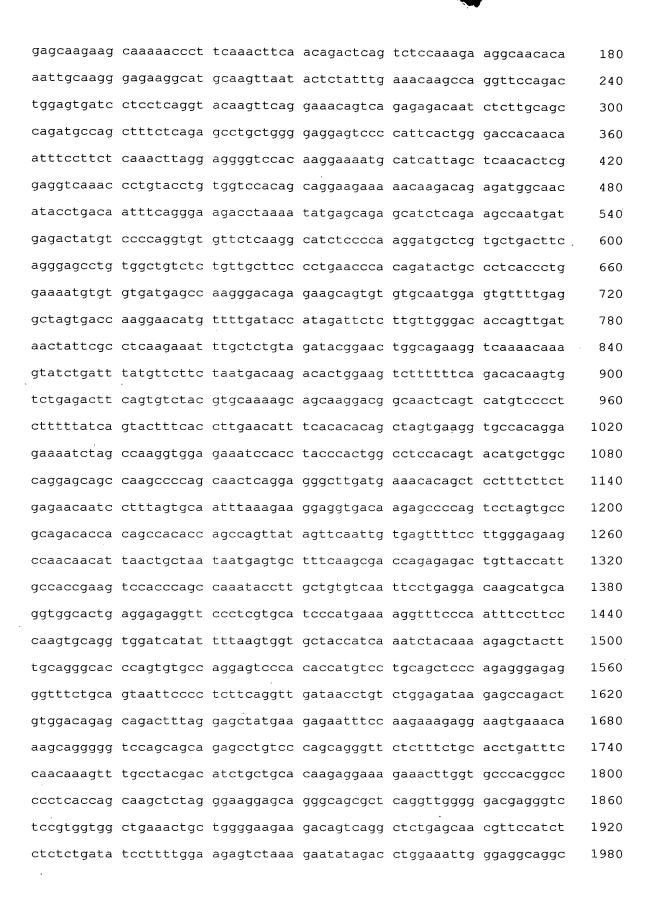
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- Glu Gly Met Gln Val Asn Thr Leu Phe Glu Thr Ser Gln Val Pro Asp 70 75 80
- Trp Ser Asp Pro Pro Gln Val Gln Val Gln Glu Thr Val Arg Glu Thr 85 90 95
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- Ser Pro Phe Thr Gly Thr Thr Thr Ile Ser Phe Ser Asn Leu Gly Gly 115° 120 125
- Val His Lys Glu Asn Ala Ser Leu Ala Gln His Ser Glu Val Lys Pro 130 135 140
- Cys Thr Cys Gly Pro Gln Gln Glu Glu Lys Gln Asp Arg Asp Gly Asn 145 150 155 160
- Ile Pro Asp Asn Phe Arg Glu Asp Leu Lys Tyr Glu Gln Ser Ile Ser 165 170 175
- Glu Ala Asn Asp Glu Thr Met Ser Pro Gly Val Phe Ser Arg His Leu 180 185 190
- Pro Lys Asp Ala Arg Ala Asp Phe Arg Glu Pro Val Ala Val Ser Val
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- Ala Ser Pro Glu Pro Thr Asp Thr Ala Leu Thr Leu Glu Asn Val Cys 210 220
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Asp Lys Thr Leu Glu Val Phe Phe Gln Thr Gln Val Ser Glu Thr Ser 290 295 300

Val Ser Thr Cys Lys Ser Ser Lys Asp Gly Asn Ser Val Met Ser Pro 305 310 315 320

Leu Phe Ile Ser Thr Phe Thr Leu Asn Ile Ser His Thr Ala Ser Glu 325 330 335

Gly Ala Thr Gly Glu Asn Leu Ala Lys Val Glu Lys Ser Thr Tyr Pro $340 \hspace{1.5cm} 345 \hspace{1.5cm} 350$

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Pro Trp Glu Lys Pro Thr Thr Leu Thr Ala Asn Asn Glu Cys Phe Gln 420 425 430

Ala Thr Arg Glu Thr Val Thr Ile Ala Thr Glu Val His Pro Ala Lys 435 440 445

Tyr Leu Ala Val Ser Ile Pro Glu Asp Lys His Ala Gly Gly Thr Glu $450 \,$ $\,$ $460 \,$

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Gln Val Gln Val Asp His Ile Leu Ser Gly Ala Thr Ile Lys Ser Thr $485 \hspace{1.5cm} 490 \hspace{1.5cm} 495$

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Arg Ser Leu Ser Ser Arg Gly Phe Ser Gln Pro Arg Leu Leu Glu Ser 755 760 765

Ser Val Asp Pro Val Asp Glu Lys Glu Leu Ser Val Thr Asp Ser Leu 770 780

Ser Ala Ala Ser Glu Thr Gly Gly Lys Glu Asn Val Asn Asn Val Ser 785 790 795 800

Gln Asp Gln Glu Lys Gln Leu Lys Met Asp His Thr Ala Phe Phe 805 810 815

Lys Lys Phe Leu Thr Cys Pro Lys Ile Leu Glu Ser Ser Val Asp Pro 820 825 830

Ile Asp Glu Ile Ser Val Ile Glu Tyr Thr Arg Ala Gly Lys Pro Glu 835 840 845

Pro Ser Glu Thr Thr Pro Gln Gly Ala Arg Glu Gly Gly Gln Ser Asn 850 855 860

Asp Gly Asn Met Gly His Glu Ala Glu Ile Gln Ser Ala Ile Leu Gln 865 870 875 880

Val Pro Cys Leu Gln Gly Thr Ile Leu Ser Glu Asn Arg Ile Ser Arg 885 890 895

Ser Gln Glu Gly Ser Met Lys Gln Glu Ala Glu Gln Ile Gln Pro Glu 900 905 910

Glu Ala Lys Thr Ala Ile Trp Gln Val Leu Gln Pro Ser Glu Gly Gly 915 920 925

Glu Arg Ile Pro Ser Gly Cys Ser Ile Gly Gln Ile Gln Glu Ser Ser 930 935 940

Asp Gly Ser Leu Gly Glu Ala Glu Gln Ser Lys Lys Asp Lys Ala Glu 945 950 955 960

Leu Ile Ser Pro Thr. Ser Pro Leu Ser Ser Cys Leu Pro Ile Met Thr 965 970 975

His Ser Ser Leu Gly Val Asp Thr His Asn Ser Thr Gly Gln Ile His 980 985 990

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Pro Ser Leu Pro Leu Glu Asn Val Gly Ser Gly Ser Arg Val Arg Glu

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Asp Leu Ala Pro Leu Ala Pro Ala Arg Pro Arg Gly Pro Leu Ile Cys 50 55 60

His Thr Gly His Glu Gln Ala Gly Arg Glu Pro Gly Pro Gly Ser Ser 65 . 70 . 75 80

Thr Lys Gly Pro Val Leu His Asp Gln Asp Thr Arg Cys Ala Phe Leu 85 90 95

Pro Arg Pro Pro Gly Pro Leu Gln Thr Arg Arg Tyr Cys Arg His Gln
100 105 110

Gly Arg Gln Gly Ser Gly Leu Gly Ala Gly Pro Gly Ala Gly Thr Trp \$115\$

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Ile Asn Arg Gly Ala Arg Gln Pro Arg Ala Gly Ala Ala Ala Gly 165 170 175

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705

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Ala Ser Ile Leu Ala Arg Asp Cys Ala Ala Ala Ala Ile Val Phe 100 105 110

Leu Val Asp Arg Phe Leu Tyr Gly Leu Asp Val Ser Gly Lys Leu Leu 115 120 125

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Pro Gln Val Val Ile Arg Gln Ala Arg Ile Ser Val Asn Ser Gly Lys 145 150 155 160

- Leu Leu Lys Ala Glu Tyr Ile Leu Ser Ser Leu Ile Ser Asn Asn Gly 165 170 175
- Ala Thr Gly Thr Trp Leu Tyr Arg Asn Glu Ser Asp Lys Val Leu Val 180 185 190
- Gln Ser Val Cys Ile Gln Ile Arg Gly Gln Ile Leu Gln Lys Leu Gly $195 \hspace{1.5cm} 200 \hspace{1.5cm} 205 \hspace{1.5cm}$
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- Phe Lys Asn Asn Pro Gln Ile Asn Leu Ser Leu Leu Lys Glu Phe Asp 260 265 270
- His His Leu Leu Ser Ala Ala Glu Ala Cys Lys Leu Ala Ala Ala Phe 275 280 285
- Ser Ala Tyr Thr Pro Leu Phe Val Leu Thr Ala Val Asn Ile Arg Gly 290 295 300
- Thr Cys Leu Leu Ser Tyr Ser Ser Ser Asn Asp Cys Pro Pro Glu Leu 305 310 315 320
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- Asp Ser Pro Trp Ser Tyr Leu Asn Ser Ser Gly Ser Ser Trp Val Ser 945 950 955 960
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